<u>Sal. abortus-equi</u> has been known as a monophasic type which is stable in both phase-1 (a) and phase-2 (e,n,x). An alternative phase is obtained by the antiserum selection for rare variations. The presence of a suppressed  $H_1$  locus in phase-2 cell is demonstrated by transduction of  $H_1^a$  allele, as well as  $H_2^{\text{enx}}$ , to the other recipient (Lederberg and Edwards, 1953). Therefore, the strain has the genotype  $H_1^aH_2^{\text{enx}}$ , but  $H_2^{\text{enx}}$  is stabilized in both activeand inactive-states. An analysis of a factor which stabilizes  $H_2$  state will be reported here.

A strain of <u>Sal. abortus-equi</u>, SW726, was used for the experiment. The strain is very slow motile in both phases. The motility is slower in phase-1 than in phase-2. Selection of a fast motile variant by NGA deep tube cultures has been unsuccessful.

For the transductional experiment, e,n,x-phase of SW726 was used as a donor and i-phase (phase-1) of diphasic <u>Sal. typhimurium</u> TW2 as a recipient. Transductional types were screened by NGA plates supplemented anti-i serum and anti e,n,x-serum at a dilution of 1/1000. Among 65 transductions obtained, 4 expressed diphasic <u>a:1,2,42 diphasic i:e,n,x</u> and remaining 19 monophasic <u>e,n,x</u>. That the hidden phase of the last type is <u>i</u> was demonstrated on three PLT22 sensitive clones by transduction to <u>Sal. paratyphi</u> B SW666 b:-. These results show that both <u>a</u> and <u>enx</u> is transduced from the phase-2 culture, and when <u>a</u> is transduced the resulted transductions remain as diphasic strains whereas when <u>e,n,x</u> is transduced some transductional clones become to be monophasic. By anti-enx NGA selection, i-phase cultures are obtained rarely from the e,n,x-monophasic transductional clones. The i-phase cultures thus obtained are also monophasic.

The stabilization of  $H_2$  state in Sal. abortus-equi is therefore caused by a gene which is linked to  $H_2$ . The controller of  $H_2$  stability will be given a symbol  $Sh_2$ . The genotype of SW726 is described as  $H_1^2H_2^{enx}$   $Sh_2^-$ .